\_\_\_\_\_\_

Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free). Reviewer: markspencer

Timestamp: [year=2009; month=3; day=12; hr=11; min=2; sec=14; ms=883; ]

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#### 

#### Reviewer Comments:

E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141>
E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151>
E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151>

<140> 10/565,646

<141> March 24, 2006

<150> PCT/FR2004/001952

<151> July 22, 2004

<150> FR/09161

<151> July 25, 2003

Please change the format of the dates above to (YYYY-MM-DD).

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<211> 648

<212> PRT

<213> Bos taurus

<400> 6

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Ser Gln Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr Ile

195 200 205

Thr Asp Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Gln Ala Leu 210 220

Asp Gly Arg Asn Lys Arg Phe Leu Arg Lys Gln Pro Leu Thr Phe Ala 225 230 235 240

"The enumeration of amino acids shall start at the first amino acid as number 1. It shall be marked below the sequence every 5 amino acids. SEQ ID # 6 amino acid number 210 is not in the correct location. Please make all necessary changes.

\*\*\*\*\*\*\*\*\*\*\*\*\*\*

# Validated By CRFValidator v 1.0.3

Application No: 10565646 Version No: 2.0

Input Set:

Output Set:

**Started:** 2009-02-05 20:45:13.750

**Finished:** 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Error code		Error Description
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	201	Mandatory field data missing in <223> in SEQ ID (1)
E	201	Mandatory field data missing in <223> in SEQ ID (1)
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E	201	Mandatory field data missing in <223> in SEQ ID (1)
E	320	Wrong Nucleic Acid Designator, ga in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gt in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (1)
E	320	Wrong Nucleic Acid Designator, ac in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gc in SEQID (1)
E	323	Invalid/missing amino acid numbering SEQID (1) POS (6764)

# Output Set:

**Started:** 2009-02-05 20:45:13.750 **Finished:** 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Err	or code	Error Description
E	320	Wrong Nucleic Acid Designator, gg in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (1)
E	320	Wrong Nucleic Acid Designator, ag in SEQID (1)
E	201	Mandatory field data missing in <223> in SEQ ID (3)
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E	201	Mandatory field data missing in $<223>$ in SEQ ID (3) This error has occured more than 20 times, will not be displayed
E	320	Wrong Nucleic Acid Designator, ga in SEQID (3)
E	320	Wrong Nucleic Acid Designator, gt in SEQID (3)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (3)
E	322	CDS location out of range SEQID (3) At Protien count (157)
E	259	Found undefined lettercode; POS (4557) SEQID(3)
E	259	Found undefined lettercode; POS (4558) SEQID(3)
E	259	Found undefined lettercode; POS (4559) SEQID(3)
E	259	Found undefined lettercode; POS (4560) SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4616, Calculated: 4620 SEQID(3)

# Output Set:

**Started:** 2009-02-05 20:45:13.750

Finished: 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Actual SeqID Count: 11

Err	or code	Error Description
E	254	The total number of bases conflicts with running total, Input: 4676, Calculated: 4680 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4734, Calculated: 4738 SEQID(3)
Ε	254	The total number of bases conflicts with running total, Input: 4782, Calculated: 4786 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4830, Calculated: 4834 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4878, Calculated: 4882 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4926, Calculated: 4930 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 4974, Calculated: 4978 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5022, Calculated: 5026 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5070, Calculated: 5074 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5118, Calculated: 5122 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5166, Calculated: 5170 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5214, Calculated: 5218 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5262, Calculated: 5266 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5310, Calculated: 5314 SEQID(3)
E	254	The total number of bases conflicts with running total, Input:

5358, Calculated: 5362 SEQID(3)

# Output Set:

**Started:** 2009-02-05 20:45:13.750

Finished: 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code	Error Description
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E 254	The total number of bases conflicts with running total, Input: 5463, Calculated: 5467 SEQID(3)
E 254	The total number of bases conflicts with running total, Input: 5523, Calculated: 5527 SEQID(3)
E 254	The total number of bases conflicts with running total, Input: 5583, Calculated: 5587 SEQID(3) This error has occured more than 20 times, will not be displayed
E 320	Wrong Nucleic Acid Designator, gc in SEQID (3)
E 322	CDS location out of range SEQID (3) At Protien count (476)
E 323	Invalid/missing amino acid numbering SEQID (3) POS (6768)
E 322	CDS location out of range SEQID (3) At Protien count (505)
E 320	Wrong Nucleic Acid Designator, gg in SEQID (3)
E 322	CDS location out of range SEQID (3) At Protien count (573)
Е 320	Wrong Nucleic Acid Designator, gg in SEQID (3)
E 322	CDS location out of range SEQID (3) At Protien count (603)
Е 320	Wrong Nucleic Acid Designator, ag in SEQID (3)
Е 322	CDS location out of range SEQID (3) At Protien count (649)
E 253	The number of bases differs from <211> Input: 8146
Е 323	Invalid/missing amino acid numbering SEQID (5) POS (2323)
Е 320	Wrong Nucleic Acid Designator, ga in SEQID (5)
Е 320	Wrong Nucleic Acid Designator, gt in SEQID (5)
Е 323	Invalid/missing amino acid numbering SEQID (5) POS (4694)
Е 320	Wrong Nucleic Acid Designator, ac in SEQID (5)

# Output Set:

**Started:** 2009-02-05 20:45:13.750 **Finished:** 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code	Error Description
E 320	Wrong Nucleic Acid Designator, gc in SEQID (5)
E 320	Wrong Nucleic Acid Designator, ag in SEQID (5) This error has occured more than 20 times, will not be displayed
W 112	Upper case found in data; Found at position(0) SeqId(6)
W 112	Upper case found in data; Found at position(3) SeqId(6)
E 259	Found undefined lettercode; POS (6) SEQID(6)
W 112	Upper case found in data; Found at position(6) SeqId(6)
E 259	Found undefined lettercode; POS (8) SEQID(6)
E 342	'n' position not defined found at POS: 9 SEQID(6)
W 112	Upper case found in data; Found at position(9) SeqId(6)
E 259	Found undefined lettercode; POS (12) SEQID(6)
W 112	Upper case found in data; Found at position(12) SeqId(6)
E 259	Found undefined lettercode; POS (15) SEQID(6)
W 112	Upper case found in data; Found at position(15) SeqId(6)
E 259	Found undefined lettercode; POS (18) SEQID(6)
W 112	Upper case found in data; Found at position(18) SeqId(6)
E 259	Found undefined lettercode; POS (20) SEQID(6)
W 112	Upper case found in data; Found at position(21) SeqId(6)
E 259	Found undefined lettercode; POS (24) SEQID(6)
W 112	Upper case found in data; Found at position(24) SeqId(6)
E 259	Found undefined lettercode; POS (26) SEQID(6)
W 112	Upper case found in data; Found at position(27) SeqId(6)
E 259	Found undefined lettercode; POS (30) SEQID(6)

# Output Set:

**Started:** 2009-02-05 20:45:13.750

Finished: 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code	Error Description
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E 259	Found undefined lettercode; POS (32) SEQID(6)
W 112	Upper case found in data; Found at position(33) SeqId(6)
E 259	Found undefined lettercode; POS (35) SEQID(6)
Е 342	'n' position not defined found at POS: 36 SEQID(6)
W 112	Upper case found in data; Found at position(36) SeqId(6)
E 259	Found undefined lettercode; POS (38) SEQID(6)
W 112	Upper case found in data; Found at position(39) SeqId(6)
E 259	Found undefined lettercode; POS (41) SEQID(6)
E 342	'n' position not defined found at POS: 42 SEQID(6)
W 112	Upper case found in data; Found at position(42) SeqId(6)
E 259	Found undefined lettercode; POS (44) SEQID(6)
W 112	Upper case found in data; Found at position(45) SeqId(6)
E 259	Found undefined lettercode; POS (47) SEQID(6)
E 259	Found undefined lettercode; POS (49) SEQID(6) This error has occured more than 20 times, will not be displayed
E 323	Invalid/missing amino acid numbering SEQID (6) POS (209)
E 323	Invalid/missing amino acid numbering SEQID (6)at Protein (210)
E 323	Invalid/missing amino acid numbering SEQID (6) POS (214)
E 323	Invalid/missing amino acid numbering SEQID (6)at Protein (215)
Е 323	Invalid/missing amino acid numbering SEQID (6) POS (219)
E 323	Invalid/missing amino acid numbering SEQID (6)at Protein (220)
Е 323	Invalid/missing amino acid numbering SEQID (6) POS (224)

# Output Set:

**Started:** 2009-02-05 20:45:13.750 **Finished:** 2009-02-05 20:47:46.221

**Elapsed:** 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code		Error Description
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (225)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (229)
E	323	Invalid/missing amino acid numbering SEQID (6) at Protein (230)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (234)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (235)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (239)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (240)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (244)
Е	323	Invalid/missing amino acid numbering SEQID (6) at Protein (245) This error has occured more than 20 times, will not be displayed
Е	331	Count of Protein differs from the <211> tag Input: 648
W	213	Artificial or Unknown found in <213> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
W	213	Artificial or Unknown found in <213> in SEQ ID (11)

#### SEQUENCE LISTING

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Lafort, Marie-Piere,
Leveziel, Hubert
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<151> July 22, 2004
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Asp Gln Tyr Trp Gln Val Leu Gly Gly Pro Val Ser	
160 165	
gga ctg agc atc ggg aca gac aag gca atg ctg ggc aca tat aac atg	4397
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met	
170 175 180	
gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc	4445
Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu	
185 190 195 200	
act cac too aut toa acc tto acc att act a ataamaata amaamaaa	4496
get cac tee agt tea gee tte ace att act g gtaaggaetg aggaggggae Ala His Ser Ser Ser Ala Phe Thr Ile Thr	4496
Ala His Ser Ser Ala Phe Thr Ile Thr	4496
	4496
Ala His Ser Ser Ala Phe Thr Ile Thr 205 210	
Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210  aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga	4556
Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210  aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga ggaaatggtg tgtaacctta caggggcaga accaggaaga tgtgggcaga gggatgtggg	4556 4616
Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210  aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga	4556

agaagetgae agaaagaaga aettatggtt eteaetttet etgaeteeaa teecag ac As	c 4734 sp
cag gtg ccc ttc tct gtg agt gtg tct cag ctg cag gcc ttg gat gga Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Gln Ala Leu Asp Gly 215 220 225	4782
agg aac aag cgc ttc ctg aga aag cag cct ctg acc ttt gcc ctc cag Arg Asn Lys Arg Phe Leu Arg Lys Gln Pro Leu Thr Phe Ala Leu Gln 230 235 240	4830
ctc cat gat ccc agt ggc tat ttg gct ggg gct gac ctt tcc tac acc Leu His Asp Pro Ser Gly Tyr Leu Ala Gly Ala Asp Leu Ser Tyr Thr 245 250 255	4878
tgg gac ttt ggt gac agt aca ggg acc ctg atc tct cgg gca ctc acg Trp Asp Phe Gly Asp Ser Thr Gly Thr Leu Ile Ser Arg Ala Leu Thr 260 265 270 275	4926
gtc act cac act tac cta gag tct ggc cca gtc act gca cag gtg gtg Val Thr His Thr Tyr Leu Glu Ser Gly Pro Val Thr Ala Gln Val Val 280 285 290	4974
ctg cag gct gcc att cct ctc acc tcc tgt ggc tcc tct cca gtt cca Leu Gln Ala Ala Ile Pro Leu Thr Ser Cys Gly Ser Ser Pro Val Pro 295 300 305	5022
ggc act aca gat agg cat gtg aca act gca gag gct cct gga acc aca Gly Thr Thr Asp Arg His Val Thr Thr Ala Glu Ala Pro Gly Thr Thr 310 315 320	5070
gct ggc caa gtg cct act aca gaa gtc atg ggc acc aca cct ggc cag Ala Gly Gln Val Pro Thr Thr Glu Val Met Gly Thr Thr Pro Gly Gln 325 330 335	5118
gtg cca act gca gag gcc cct ggc acc aca gtt ggg tgg gtg cca acc Val Pro Thr Ala Glu Ala Pro Gly Thr Thr Val Gly Trp Val Pro Thr 340 345 350 355	5166
aca gag gat gta ggt acc aca cct gag cag gtg gca acc tcc aaa gtc  Thr Glu Asp Val Gly Thr Thr Pro Glu Gln Val Ala Thr Ser Lys Val  360 365 370	5214
tta agt aca aca cca gtg gag atg cca act gca aaa gct aca ggt agg Leu Ser Thr Thr Pro Val Glu Met Pro Thr Ala Lys Ala Thr Gly Arg 375 380 385	5262
aca cct gaa gtg tca act aca gag ccc tct gga acc aca gtt aca cag Thr Pro Glu Val Ser Thr Thr Glu Pro Ser Gly Thr Thr Val Thr Gln 390 395 400	5310
gga aca act cca gag ctg gtg gag acc aca gct gga gag gtg tcc act Gly Thr Thr Pro Glu Leu Val Glu Thr Thr Ala Gly Glu Val Ser Thr 405 410 415	5358

cct gag cct gcg ggt tca aat act agc tca ttc atg cct aca gaa ggt Pro Glu Pro Ala Gly Ser Asn Thr Ser Ser Phe Met Pro Thr Glu Gly 420 425 430 430	5406
act gca g gtaagggggc caccatgaat gagttcatag aggtggggca tttgtcacag Thr Ala	5463
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Leu Pro Lys Glu Ala 510

tgc atg gac atc tca tcg cca ggg tgt cag ctg cct gcc cag cgg ctg Cys Met Asp Ile Ser Ser Pro Gly Cys Gln Leu Pro Ala Gln Arg Leu 515 520 525	6938
tgt cag cct gtg ccc ccc age cca gcc tgc cag ctg gtt ttg cac cag Cys Gln Pro Val Pro Pro Ser Pro Ala Cys Gln Leu Val Leu His Gln 530 535 540	6986
gta ctg aag ggt ggc tca ggg acc tac tgc ctc aat gtg tct ttg gct Val Leu Lys Gly Gly Ser Gly Thr Tyr Cys Leu Asn Val Ser Leu Ala 545 550 555	7034
gat gcc aat agc ctg gcg atg gtc agc acc cag ctt gtc atg cct g Asp Ala Asn Ser Leu Ala Met Val Ser Thr Gln Leu Val Met Pro 560 565 570	7080
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gca ggc ctc agg cag gct cct ctg ttc gtg ggc atc ttg ctg gtg cta Ala Gly Leu Arg Gln Ala Pro Leu Phe Val Gly Ile Leu Leu Val Leu 580 585 590	7243
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